# Class 04: Linear Regression and Regularization

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# Recap: Empirical Risk Minimization (ERM)

#### Setup:

- sample  $\mathcal{D}_n = \{(X_i, Y_i)\}_{i=1}^n$ ,
- a loss function L,
- a function class .F

#### Empirical risk minimization:

$$\min_{f \in \mathcal{F}} R_{\mathsf{emp}}(f) = \min_{f \in \mathcal{F}} \frac{1}{n} \sum_{i=1}^{n} L(Y_i, f(X_i)).$$

Let  $\widehat{f}$  denote the empirical risk minimizer. We have seen that  $R_{\rm emp}(\widehat{f})$  can be a poor proxy for the generalization error

$$\mathbf{E}_{X,Y}[L(Y,\widehat{f}(X))|\widehat{f}].$$

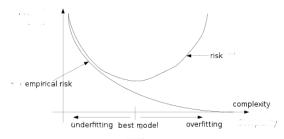
# Overfitting and Underfitting

### Overfitting:

By choosing  $\mathcal F$  "large enough", it easy to achieve a small empirical risk. However, the empirical risk minimizer typically does not generalize well to future data  $(X,Y)\sim P$ .

### **Underfitting**:

On the other hand, if  $\mathcal{F}$  is too small, we may be unable to capture important characteristics of the problem we want to learn.



### Structural Risk Minimization

What is good a practical strategy for controlling the capacity of  $\mathcal{F}$ ?

**Structural Risk Minimization**: Working with a nested sequence of increasingly complex function classes

$$\mathcal{F}_1 \subset \mathcal{F}_2 \subset \ldots \subset \mathcal{F}_N$$
,

perform ERM separately for each class, and then pick the empirical risk minimizer from the least complex class with reasonable empirical error (i.e. within reach of more complex classes).

The underlying principle is known as **Occam's Razor** (William Ockham, 1287–1347) or **Principle of Parsimony**:

Among all models that explain the data well, take the simplest one.

### Structural Risk Minimization

The idea of Structural Risk Minimization (SRM) is commonly used for model selection in regression models, often combined with criteria such as Akaike's information criterion (AIC) or Schwartz' Bayesian information criterion (BIC).

A problem with SRM is that it tends to be computationally inefficient:

- we have to compute the ERM for each function class under consideration
- the number of candidate function classes can be prohibitively large

## Regularization

A conceptually closely related strategy is **Regularization**.

- ullet we work with a large a function class  ${\mathcal F}$ ,
- we add an additional term (the regularizer) to the empirical risk,
- the purpose of the regularizer is to penalize more complex hypotheses in  $\mathcal{F}$ .

This yields regularized empirical risk minimization:

$$\min_{f \in \mathcal{F}} R_{\mathsf{emp}}(f) + \lambda \Omega(f),$$

where

- $\Omega: \mathcal{F} \to \mathbb{R}_+$  is called regularizer,
- $\lambda \ge 0$  is called **regularization parameter**.

# Regularized empirical risk minimization

$$\min_{f \in \mathcal{F}} R_{\mathsf{emp}}(f) + \lambda \Omega(f),$$

The regularization parameter controls the trade-off between

- achieving low empirical risk  $(\lambda \to 0)$ ,
- controlling model complexity  $(\lambda \to \infty)$ .

From an optimization standpoint,  $\lambda$  can be interpreted as a Lagrangian multiplier for the constrained optimization problem

$$\min_{f \in \mathcal{F}} R_{\mathsf{emp}}(f)$$
 subject to  $\Omega(f) \leq r,$ 

## Regularized empirical risk minimization

Many popular machine learning algorithms can be put into the framework of regularized empirical risk minimization.

They differ by the choice of the loss L and the regularizer  $\Omega$ .

#### Examples:

- Ridge Regression,
- Support Vector Classification,
- Support Vector Regression,
- Logistic Regression,
- Lasso, Elastic Net, Group Lasso

We want to learn a function that predicts Y from X.

More specifically, our target is

$$f_L^*(x) = \mathbf{E}[Y|X=x], \ x \in \mathcal{X},$$

i.e. the corresponding loss function is squared loss

$$L(Y, f(X)) = (Y - f(X))^2.$$

The function class  $\mathcal F$  is given by

$$\mathcal{F} = \left\{ f: \ x \mapsto f(x) = \sum_{j=1}^{D} w_j \phi_j(x), \ w \in \mathbb{R}^D \right\},\,$$

where the  $\{\phi_j\}_{j=1}^D$  are fixed functions  $\mathcal{X} \to \mathbb{R}$ .

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where the  $\{\phi_j\}_{j=1}^D$  are fixed functions  $\mathcal{X} \to \mathbb{R}$ .

The use of a input transformation (or feature map)

$$\mathcal{X} \to \mathbb{R}^D$$

$$x \mapsto \Phi(x) := \begin{pmatrix} \phi_1(x) \\ \vdots \\ \phi_D(x) \end{pmatrix}$$

allows for additional modeling flexibility.

#### Examples of Input transformations:

(1) No transformation: keep the original features, and only these

$$\phi_1(x) = x_1, \dots, \phi_D(x) = x_d, \quad D = d.$$

(2) Interactions up to order two

$$\phi_1(x) = x_1, \dots, \phi_d(x) = x_d,$$
  

$$\phi_{d+1}(x) = x_1 x_2, \dots, \phi_D = x_{d-1} x_d, \quad D = d(d+1)/2.$$

(3) Polynomial terms  $(\mathcal{X} \subseteq \mathbb{R})$ 

$$\phi_1(x) = 1, \ \phi_2(x) = x, \dots, \ \phi_D(x) = x^{D-1}, \phi_D(x) = x^D.$$

(4) Trigonometric polynomials  $(\mathcal{X} \subseteq [0,1])$ 

$$\phi_1(x) = 1, \ \phi_2(x) = \sin(2\pi x), \phi_3(x) = \cos(2\pi x), \dots,$$
$$\phi_{D-1}(x) = \sin(2\pi N x), \phi_D(x) = \cos(2\pi N x).$$

for some positive integer N.

many more possibilities: splines, wavelets, fractional polynomials,

Note: linear regression does not exclude nonlinear transformations of the inputs.

Linear Regression because

$$f(x) = \sum_{j=1}^{D} \phi_j(x) w_j = \Phi(x)^{\top} w, \quad w = \begin{pmatrix} w_1 \\ \vdots \\ w_D \end{pmatrix}$$

is a linear function in w.

With  $L(Y, f(X)) = \{Y - f(X)\}^2$ , empirical risk minimization amounts to least squares estimation.

Denote by

$$\mathbf{\Phi} = \begin{pmatrix} \Phi(X_1)^{\top} \\ \vdots \\ \Phi(X_n)^{\top} \end{pmatrix} = \begin{pmatrix} \phi_1(X_1) & \dots & \phi_D(X_1) \\ \vdots & \dots & \vdots \\ \phi_1(X_n) & \dots & \phi_D(X_n). \end{pmatrix}$$

the  $\mathbb{R}^{n \times D}$  feature matrix or design matrix, and let

$$\mathbf{y} = \begin{pmatrix} Y_1 \\ \vdots \\ Y_n \end{pmatrix}$$

be the vector of outputs.

Unless stated explicitly, there is no intercept in the model – i.e. the matrix  $\Phi$  does not contain a column of ones.

Note that we do not need an intercept whenever y and all columns of  $\Phi$  are centered (i.e., they sum up to zero).

$$\min_{f \in \mathcal{F}} R_{\mathsf{emp}}(f) = \min_{f \in \mathcal{F}} \frac{1}{n} \sum_{i=1}^{n} L(Y_i, f(X_i))$$

$$= \min_{w \in \mathbb{R}^D} \frac{1}{n} \sum_{i=1}^{n} (Y_i - \Phi(X_i)^{\top} w)^2$$

$$= \min_{w \in \mathbb{R}^D} \frac{1}{n} ||\mathbf{y} - \mathbf{\Phi} w||_2^2$$

As in Homework 0, it can be shown any  $\widehat{w}_{\mathrm{LS}}$  satisfying the *normal equations* 

$$\frac{1}{n} \mathbf{\Phi}^{\top} \mathbf{\Phi} \widehat{w}_{\mathsf{LS}} = \frac{1}{n} \mathbf{\Phi}^{\top} \mathbf{y}$$

is optimal.  $\widehat{w}_{LS}$  is called a least squares estimator. It is unique whenever  $\Phi$  is non-singular  $\Leftrightarrow \Phi^{\top}\Phi$  is invertible. In this case,

$$\widehat{w}_{\mathsf{LS}} = (\mathbf{\Phi}^{\top} \mathbf{\Phi})^{-1} \mathbf{\Phi}^{\top} \mathbf{y}.$$

Is least squares estimation a good idea?

Least squares is prone to overfit whenever  ${\cal D}$  becomes large relative to n.

In the scenario  $D \geq n$ , it is possible that  $\mathrm{range}(\Phi) = \mathbb{R}^n$ , where  $\mathrm{range}(\Phi)$  denotes the column space of  $\Phi$ . In this situation,

$$\Phi \widehat{w}_{LS} = \mathbf{y},$$

i.e., we perfectly fit the observed sample, but likely do poorly on unseen data.

Example: Pima Indian Diabetes data set.

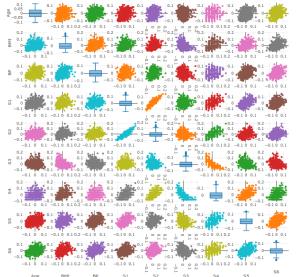
n=442 diabetes patients

Target variable to be predicted is a measure of disease progression.

Covariates (d = 10):

- age
- sex
- body mass index
- average blood pressure
- six blood serum measurements

Scatterplot matrix of continuous variables (all except for sex):



### Correlation matrix:

$\lceil \circ \rceil$	0.17	0.19	0.34	0.26	0.22	-0.08	0.2	0.27	0.3
0	0	0.09	0.24	0.04	0.14	-0.38	0.33	0.15	0.21
0	0	0	0.4	0.25	0.26	-0.37	0.41	0.45	0.39
0	0	0	0	0.24	0.19	-0.18	0.26	0.39	0.39
0	0	0	0	0	0.9	0.05	0.54	0.52	0.33
0	0	0	0	0	0	-0.2	0.66	0.32	0.29
0	0	0	0	0	0	0	-0.74	-0.4	-0.27
0	0	0	0	0	0	0	0	0.62	0.42
0	0	0	0	0	0	0	0	0	0.46
0	0	0	0	0	0	0	0	0	0

We fit a linear regression model using all d=10 covariates.

Training error	Leave-One-Out Error		
2,860	2,988		

To verify that least squares tends to overfit we randomly permute the rows of  $\Phi$  and append them as additional columns. This is repeated multiple times:

#permutations	D	Training error	Leave-One-Out Error
1	20	2,816	3,097
2	30	2,739	3,146
3	40	2,618	3,172
4	50	2,561	3,257

This could be repeated by appending as many permuted versions of  $\Phi$  until the training error drops to zero.

The idea behind the permutations is that they break the correlation to the response y; hence the extra columns take the role of "noise variables".

In the following, we present a simplified analysis of least squares, that still conveys the main insights.

(I) Only the Y's are random, i.e

$$\mathcal{D}_n = \{(x_1, Y_1), \dots, (x_n, Y_n)\},\$$

where the  $\{x_i\}_{i=1}^n$  are considered fixed.

(II) Additive and homoskedastic noise

$$Y_i = f^*(x_i) + \varepsilon_i, \quad i = 1, \dots, n,$$

where for the noise terms we assume that

- $\mathbf{E}[\varepsilon_i] = 0$ ,
- $\mathbf{E}[\varepsilon_i^2] = \sigma^2$ , i = 1, ..., n,
- The  $\{\varepsilon_i\}_{i=1}^n$  are uncorrelated.

Suppose hypothetically we are also given a second data set, say, for the purpose of validation:

$$\mathcal{D}'_n = \{(x_1, Y'_1), \dots, (x_n, Y'_n)\},\$$

where

$$Y_i' = f^*(x_i) + \varepsilon_i',$$

and  $\varepsilon_i'$  has the same distribution as  $\varepsilon_i$ ,  $i=1,\ldots,n$ .

 $\mathcal{D}_n$ :

features	"ground truth"	noise	response
$x_1$	$f^*(x_1)$	$arepsilon_1$	$Y_1$
:	:	:	:
$x_n$	$f^*(x_n)$	$arepsilon_n$	$Y_n$

$$\mathcal{D}'_n$$
:

features	"ground truth"	noise	response
$x_1$	$f^*(x_1)$	$arepsilon_1'$	$Y_1'$
÷	:	:	:
$x_n$	$f^*(x_n)$	$arepsilon_n'$	$Y_n'$

# Least Squares Estimation

The statistical performance of  $w \in \mathbb{R}^D$  can be quantified by

$$MSE(w) = \mathbf{E}\left[\frac{1}{n}\sum_{i=1}^{n}(Y_i' - \Phi(x_i)^{\top}w)^2\right]$$

where  $\mathbf{E}[\cdot]$  denotes the expectation with respect to  $\varepsilon'$ .

Clearly, the optimal MSE is given by

$$\mathsf{MSE}(w^*) = \min_{w \in \mathbb{R}^D} \mathbf{E} \left[ \frac{1}{n} \sum_{i=1}^n (Y_i' - \Phi(x_i)^\top w)^2 \right]$$

This our benchmark, and we will compare how several estimators perform relative to this benchmark.

It is not hard to show that

$$\begin{aligned} \mathsf{MSE}(w^*) &= \min_{w \in \mathbb{R}^D} \mathbf{E} \left[ \frac{1}{n} \sum_{i=1}^n (Y_i' - \Phi(x_i)^\top w)^2 \right] \\ &= \underbrace{\sigma^2}_{\mathsf{noise}} + \underbrace{\min_{w \in \mathbb{R}^D} \frac{1}{n} \| \boldsymbol{f}^* - \boldsymbol{\Phi} w \|_2^2}_{\mathsf{approximation error}}, \end{aligned}$$

where

$$\mathbf{f}^* = (f^*(x_1) \quad \dots \quad f^*(x_n))^\top.$$

## Least Squares Estimation

One can show the following for the least squares estimator:

$$\begin{split} \mathbf{E}[\mathsf{MSE}(\widehat{w}_{\mathsf{LS}})] &= \underbrace{\sigma^2}_{\mathsf{noise}} + \underbrace{\min_{w \in \mathbb{R}^D} \frac{1}{n} \| \boldsymbol{f}^* - \boldsymbol{\Phi} w \|_2^2}_{\mathsf{approximation \ error}} + \underbrace{\frac{\sigma^2 \mathsf{rank}(\boldsymbol{\Phi})}{n}}_{\mathsf{estimation \ error}} \end{split}$$

Note that since  $\widehat{w}_{LS}$  is random, so is  $MSE(\widehat{w}_{LS})$  – that's why we take the expectation  $\mathbf{E}[\cdot]$  w.r.t.  $\varepsilon$  from the data set  $\mathcal{D}_n$  from which  $\widehat{w}_{LS}$  was obtained.

# Least Squares Estimation

$$\mathbf{E}[\mathsf{MSE}(\widehat{w}_{\mathsf{LS}})] = \mathsf{MSE}(w^*) + \frac{\sigma^2\mathsf{rank}(\boldsymbol{\Phi})}{n}$$

Note that typically

$$\mathsf{rank}(\mathbf{\Phi}) = \min\{D, n\}.$$

In this case, the result implies that least squares is not even statistically consistent unless D is small relative to n.

Least squares can be a poor method even if n > D (and is useless if D > n).

A variety of properties of least squares regression and several other methods can be better understood in terms of the **singular value** decomposition (SVD) of  $\Phi$ .

Without any additional assumption on  $\Phi$ , we have

$$\boldsymbol{\Phi} = \mathbf{U}\mathbf{S}\mathbf{V}^\top$$

- $\bullet$   $\mathbf{U} \in \mathbb{R}^{n \times n}$  such that  $\mathbf{U}^{\top} \mathbf{U} = \mathbf{U} \mathbf{U}^{\top} = I_n$ ,
- $\bullet$   $\mathbf{S} \in \mathbb{R}^{n \times D}$  and  $m = \min\{n, D\}$  such that

$$\mathbf{S} = \begin{bmatrix} \mathsf{diag}(s_1, \dots, s_m) & \mathbf{0}_{m \times (D-m)} \\ \mathbf{0}_{(n-m) \times m} & \mathbf{0}_{(n-m) \times (D-m)} \end{bmatrix},$$

 $\bullet \ \mathbf{V} \in \mathbb{R}^{D \times D} \ \text{such that} \ \mathbf{V}^\top \mathbf{V} = \mathbf{V} \mathbf{V}^\top = I_D.$ 

$$\mathbf{\Phi} = \mathbf{U}\mathbf{S}\mathbf{V}^{ op}$$

- $\bullet$   $\mathbf{U} \in \mathbb{R}^{n \times n}$  such that  $\mathbf{U}^{\top} \mathbf{U} = I_n$ ,
- $\bullet$   $\mathbf{S} \in \mathbb{R}^{n \times D}$  and  $m = \min\{n, D\}$  such that

$$\mathbf{S} = \begin{bmatrix} \mathsf{diag}(s_1, \dots, s_m) & \mathbf{0}_{m \times (D-m)} \\ \mathbf{0}_{(n-m) \times m} & \mathbf{0}_{(n-m) \times (D-m)} \end{bmatrix},$$

•  $\mathbf{V} \in \mathbb{R}^{D \times D}$  such that  $\mathbf{V}^{\top} \mathbf{V} = I_D$ .

Without loss of generality, we assume that the singular values

$$s_1 \geq \ldots \geq s_m \geq 0$$

are ordered.

"Economy" SVD (default in R, optional in python and MATLAB) (recall  $m = \min\{n, D\}$ ):

- ullet  $\mathbf{U} \in \mathbb{R}^{n \times m}$  such that  $\mathbf{U}^{\top}\mathbf{U} = I_m$ ,
- $\mathbf{S} = \mathsf{diag}(s_1, \ldots, s_m)$ ,
- $\mathbf{V} \in \mathbb{R}^{D \times m}$  such that  $\mathbf{V}^{\top} \mathbf{V} = I_m$ .

If 
$$n \leq D$$
:  $\mathbf{U}\mathbf{U}^{\top} = I_n$ ,

If 
$$n \geq D$$
:  $\mathbf{V}\mathbf{V}^{\top} = I_D$ .

### Basic properties:

- ullet The columns of U are called **left singular values**
- ullet The number of non-zero singular values equals  $\mathrm{rank}(oldsymbol{\Phi}) \leq m$
- ullet The columns of V are called **right singular values**

Moreover, we have

$$oldsymbol{\Phi}^{ op}oldsymbol{\Phi} = \mathbf{V}\mathbf{S}^2\mathbf{V}^{ op} = \mathbf{V}oldsymbol{\Lambda}\mathbf{V}^{ op}, \quad oldsymbol{\Lambda} = \mathsf{diag}(\underbrace{s_1^2}_{\lambda_1}, \ldots, \underbrace{s_m^2}_{\lambda_m})$$

i.e., the columns of V are the eigenvectors and  $\lambda_1 \geq \ldots \geq \lambda_m \geq 0$  are the eigenvalues of  $\Phi^{\top}\Phi$ .

Finally, it can be shown that

$$\mathbf{\Phi}\widehat{w}_{\mathsf{LS}} = \mathbf{U}\mathbf{U}^{\mathsf{T}}\mathbf{y},$$

i.e.,  $\mathbf{U}\mathbf{U}^{\top}$  equals the "Hat matrix" of the least squares fit.

Let  $\|M\|_F = \sqrt{\sum_{i,j} M_{ij}^2}$  denote the Frobenius norm of a matrix M.

#### Eckart-Young Theorem (1936)

For  $r \leq m$ , consider the minimization problem

$$\min_{\mathbf{Z}: \ \mathsf{rank}(\mathbf{Z}) \leq r} \|\mathbf{\Phi} - \mathbf{Z}\|_F^2$$
 .

Then an optimal solution for Z is given by

$$\mathbf{\Phi}_r = \mathbf{U}_r \mathbf{S}_r \mathbf{V}_r^{ op},$$

where  $\mathbf{U}_r$  and  $\mathbf{V}_r^{\top}$  contain the r columns of  $\mathbf{U}$  respectively rows of  $\mathbf{V}^{\top}$  corresponding to the largest r singular values, and

$$\mathbf{S}_r = \mathsf{diag}(s_1, \dots, s_r).$$

The matrix  $\Phi_r = \mathbf{U}_r \mathbf{S}_r \mathbf{V}_r^{\top}$  is called the best rank-r approximation to  $\Phi$ .

It is not hard to show that

$$\|\mathbf{\Phi} - \mathbf{\Phi}_r\|_F^2 = \sum_{i: i > r} s_j^2,$$

and thus

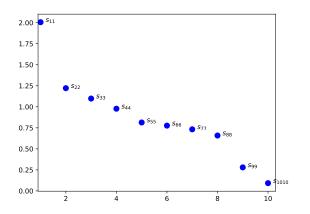
$$\mathtt{range}(\mathbf{\Phi}_r) pprox \mathtt{range}(\mathbf{\Phi})$$

if the term

$$\sum_{j:\,j>r} s_j^2$$

is small.

Plot of the sequence of singular values for the diabetes data set:



Such plot is often called "Screeplot".

One way of improving over standard least squares is to work with  $\Phi_r$  respectively  $\mathbf{U}_r$  (we have  $\mathtt{range}(\Phi_r) = \mathtt{range}(\mathbf{U}_r)$ ) instead of  $\Phi$ .

Denote by  $\widehat{w}_r$  the least squares estimator corresponding to  $\Phi_r$ .

One can show that

$$\mathbf{E}[\mathsf{MSE}(\widehat{w}_r)] = \mathsf{MSE}(w^*) + \frac{1}{n} \sum_{j > r} s_j^2 \{\alpha_j^*\}^2 + \frac{\sigma^2 r}{n}, \quad \alpha^* := V^\top w^*.$$

Hence if  $r \ll D$  and  $\sum_{j>r} s_j^2$  is small, the MSE gets substantially reduced.

Suppose D>n. Then the least squares estimator defined by the normal equations

$$\frac{1}{n} \mathbf{\Phi}^{\top} \mathbf{\Phi} \widehat{w}_{\mathsf{LS}} = \frac{1}{n} \mathbf{\Phi}^{\top} \mathbf{y}$$

is not unique.

One way of enforcing a unique solution is by adding multiple of the identity  $\lambda I_D$ ,  $\lambda > 0$ , to  $\Phi^{\top}\Phi$ : this yields

$$\begin{split} & \left(\frac{1}{n} \mathbf{\Phi}^{\top} \mathbf{\Phi} + \lambda I_{D}\right) \widehat{w}_{\mathsf{ridge}} = \frac{1}{n} \mathbf{\Phi}^{\top} \mathbf{y} \\ & \Rightarrow \ \widehat{w}_{\mathsf{ridge}} = \left(\frac{1}{n} \mathbf{\Phi}^{\top} \mathbf{\Phi} + \lambda I_{D}\right)^{-1} \frac{1}{n} \mathbf{\Phi}^{\top} \mathbf{y} \end{split}$$

Equivalently, we can define the ridge estimator by

$$\widehat{w}_{\mathsf{ridge}} = \operatorname*{argmin}_{w \in \mathbb{R}^D} \frac{1}{n} \|\mathbf{y} - \mathbf{\Phi} w\|_2^2 + \lambda \|w\|_2^2$$

which is equivalent to finding

$$\underset{f \in \mathcal{F}}{\operatorname{argmin}} R_{\mathsf{emp}}(f) + \lambda \Omega(f),$$

where

- $\bullet$   $R_{\text{emp}}$  is w.r.t. squared loss,
- ullet  ${\cal F}$  is the linear class on slide 9,

• 
$$\Omega(f) = \Omega\left(\sum_{j=1}^{D} w_j \phi_j\right) = ||w(f)||_2^2$$

It can be shown that the ridge regression fit can be expressed as

$$\mathbf{\Phi} \widehat{w}_{\mathsf{ridge}} = \mathbf{U} \; \mathsf{diag} \left( \frac{\lambda_1}{\lambda_1 + \lambda}, \dots, \frac{\lambda_m}{\lambda_m + \lambda} \right) \; \mathbf{U}^{\top} \mathbf{y},$$

recalling that  $\lambda_j = s_j^2$ ,  $j = 1, \dots, m$ .

This is to be compared to

$$\mathbf{\Phi}\widehat{w}_{\mathsf{LS}} = \mathbf{U}\mathbf{U}^{\mathsf{T}}\mathbf{y},$$

We can think of each of the columns of U as one specific direction among m directions associated with range( $\Phi$ ).

Loosely speaking, the corresponding eigenvalues  $\{\lambda_j\}_{j=1}^m$  represent the "prominence" of these directions in  $\mathtt{range}(\Phi)$ .

The larger  $\lambda_j$ , the more "prominent" the corresponding direction.

$$\mathbf{\Phi} \widehat{w}_{\mathsf{ridge}} = \mathbf{U} \; \mathsf{diag} \left( \frac{\lambda_1}{\lambda_1 + \lambda}, \dots, \frac{\lambda_m}{\lambda_m + \lambda} \right) \; \mathbf{U}^\top \mathbf{y},$$

We may intepret

$$\frac{\lambda_j}{\lambda_j + \lambda}$$

as a weight assigned to direction j, j = 1, ..., m.

The presence of  $\lambda$  in the denominator dampens the influence of directions with small  $\lambda_j$  on the regression fit much more strongly.

It can be shown that

$$\mathbf{E}[\mathsf{MSE}(\widehat{w}_{\mathsf{ridge}})] = \mathsf{MSE}(w^*) + \sum_{j=1}^D \{\alpha_j^*\}^2 \lambda_j \left(\frac{\lambda}{\lambda + \lambda_j}\right)^2, \quad \boldsymbol{\alpha}^* := \boldsymbol{V}^\top \boldsymbol{w}^*,$$

$$+\frac{\sigma^2}{n}\sum_{j=1}^D \left(\frac{\lambda_j}{\lambda_j+\lambda}\right)^2,$$

•

Ridge Regression works well if the first few eigenvalues, say  $\lambda_1, \ldots, \lambda_r$ , are dominant, while the rest  $\lambda_{r+1}, \ldots, \lambda_D$  are small, and  $\lambda$  is chosen such that

$$\lambda_{r+1} \ll \lambda \ll \lambda_r$$
.

$$\mathbf{E}[\mathsf{MSE}(\widehat{w}_{\mathsf{ridge}})] = \mathsf{MSE}(w^*) + \sum_{j=1}^{D} \{\alpha_j^*\}^2 \lambda_j \left(\frac{\lambda}{\lambda + \lambda_j}\right)^2, \quad \boldsymbol{\alpha}^* := \boldsymbol{V}^\top \boldsymbol{w}^*,$$

$$+\frac{\sigma^2}{n}\sum_{j=1}^D \left(\frac{\lambda_j}{\lambda_j + \lambda}\right)^2$$

•

Ridge Regression does not work well if the magnitude of the  $\alpha_j^*$ 's is not well aligned with the size of the  $\lambda_j$  (i.e. large  $\alpha_j^*$  for small  $\lambda_j$ ).

•

In particular, ridge regression tends to be suboptimal for **sparse**  $w^*$ . Ridge regression does not promote a parsimonious model.

Example: Pima Indian Diabetes data set (continued).

n=442 diabetes patients

Target variable to be predicted is a measure of disease progression.

Covariates (d = 10):

- age
- sex
- body mass index
- average blood pressure
- six blood serum measurements

This time, we fit a "quadratic model", i.e. in addition to the original 10 covariates, we consider quadratic terms (except for sex)

$$X_1^2, X_3^2, \dots, X_d^2,$$

and all first-order interactions

$$X_1X_2, X_1X_3, \dots, X_{d-1}X_d.$$

This yields  $D = 10 + 9 + \binom{10}{2} = 64$ .

D=64 is fairly large number of features relative to only n=442 samples. Least squares cannot be expected to work well.

Important "pre-processing" steps before applying ridge regression:

**1** Center y and the columns of  $\Phi$ :

$$y_i \leftarrow y_i - \frac{1}{n} \sum_{i'=1}^n y_{i'}, \qquad \mathbf{\Phi}_{ij} \leftarrow \mathbf{\Phi}_{ij} - \frac{1}{n} \sum_{i'=1}^n \mathbf{\Phi}_{i'j},$$

$$i = 1, \dots, n, j = 1, \dots, D.$$

2 Scale the columns of  $\Phi$  so that their norms are identical:

$$\Phi_{\bullet j} \leftarrow \Phi_{\bullet j} / \|\Phi_{\bullet j}\|_2, \ j = 1, \dots, D.$$

On the importance of column normalization when using ridge regression:

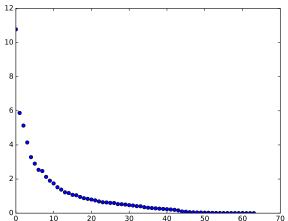
If the features are not on the same scale, we essentially penalize each feature differently. This is typically not appropriate.

Let 
$$S = \operatorname{diag}(\|\mathbf{\Phi}_{\bullet 1}\|_2, \dots, \|\mathbf{\Phi}_{\bullet D}\|_2)$$
. Then:

$$\begin{split} \min_{w \in \mathbb{R}^D} \frac{1}{n} \|\mathbf{y} - \mathbf{\Phi} w\|_2^2 + \lambda \|w\|_2^2 &= \min_{w \in \mathbb{R}^D} \frac{1}{n} \|\mathbf{y} - (\mathbf{\Phi} S^{-1})(Sw)\|_2^2 + \lambda \|w\|_2^2 \\ &= \min_{\theta \in \mathbb{R}^D} \frac{1}{n} \|\mathbf{y} - (\mathbf{\Phi} S^{-1})\theta\|_2^2 + \lambda \|S^{-1}\theta\|_2^2 \\ &= \min_{\theta \in \mathbb{R}^D} \frac{1}{n} \|\mathbf{y} - (\mathbf{\Phi} S^{-1})\theta\|_2^2 + \lambda \sum_{i=1}^D \frac{w_i^2}{\|\mathbf{\Phi}_{\bullet j}\|_2^2}. \end{split}$$

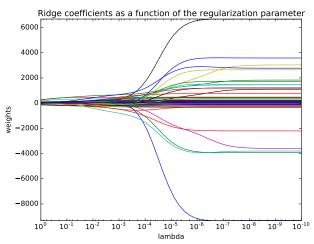
 $\rightsquigarrow$  features with small norm (scale) are penalized more strongly.

Back to the data set. "Scree plot" of the eigenvalues of  $\Phi^{\top}\Phi$  (after centering / scaling):



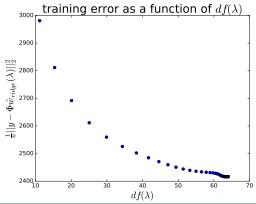
4 prominent eigenvalues, several ( $\sim 10$ ) very small eigenvalues.

Ridge coefficients (or weights)  $\widehat{w}_{\mathsf{ridge},j}$ ,  $j=1,\ldots,D$ , as a function of the regularization parameter:



"Effective #of variables" / "degrees of freedom" in ridge regression:

$$\mathrm{df}(\lambda) = \sum_{j=1}^{D} \left(\frac{\lambda_j}{\lambda_j + \lambda}\right)^2$$



Another regularization method that works better in a "sparse regime" is the lasso:

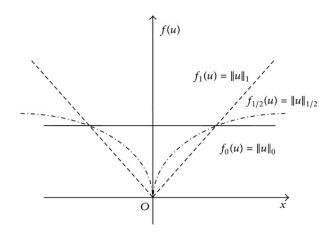
$$\widehat{w}_{\mathsf{lasso}} \in \operatorname*{argmin}_{w \in \mathbb{R}^D} \frac{1}{2n} \|\mathbf{y} - \mathbf{\Phi} w\|_2^2 + \lambda \|w\|_1.$$

It can be motivated as a convex relaxation of

$$\widehat{w}_{\ell_0} \in \underset{w \in \mathbb{R}^D}{\operatorname{argmin}} \frac{1}{2n} \|\mathbf{y} - \mathbf{\Phi}w\|_2^2 + \lambda \|w\|_0,$$
$$\|w\|_0 := \sum_{j=1}^D I(w_j \neq 0)$$

which is computationally intractable.

 $\|\cdot\|_1$  is the tightest convex approximation to  $\|\cdot\|_0$  on the cube  $B^d_\infty$ :



The  $\ell_1$ -norm regularizer produces sparse solutions, i.e., depending on  $\lambda$ , many entries of  $\widehat{w}_{\text{lasso}}$  are exactly zero.

That is: the lasso implicitly performs feature selection.

In the case that  $\frac{1}{n}\mathbf{\Phi}^{\top}\mathbf{\Phi} = \mathsf{diag}(\lambda_1,\ldots,\lambda_D)$ , it can be shown that

$$\widehat{w}_{\mathsf{lasso},j} = \frac{\operatorname{sign}\left(\frac{1}{n}\sum_{i=1}^{n}\Phi_{j}(X_{i})Y_{i}\right)\cdot \max\{\left|\frac{1}{n}\sum_{i=1}^{n}\Phi_{j}(X_{i})Y_{i}\right| - \lambda, 0\}}{\lambda_{j}},$$

$$j = 1, \dots, D.$$

This is known as "Soft Thresholding" (Donoho and Johnstone, 1994).

In the case that  $\frac{1}{n}\mathbf{\Phi}^{\top}\mathbf{\Phi}=\mathsf{diag}(\lambda_1,\ldots,\lambda_D)$ , it can be shown that

$$\widehat{w}_{\mathsf{lasso},j} = \frac{\mathsf{sign}\left(\frac{1}{n}\sum_{i=1}^{n}\Phi_{j}(X_{i})Y_{i}\right) \cdot \max\{\left|\frac{1}{n}\sum_{i=1}^{n}\Phi_{j}(X_{i})Y_{i}\right| - \lambda, 0\}}{\lambda_{j}},$$

 $j=1,\ldots,D.$ 

In particular, this implies that

$$\left| \frac{1}{n} \sum_{i=1}^{n} \Phi_{j}(X_{i}) Y_{i} \right| < \lambda \quad \Longrightarrow \ \widehat{w}_{\mathsf{lasso}, j} = 0, \ \ j = 1, \dots, D.$$

Hard thresholding (Left) vs. Soft Thresholding (Right)

